

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/559,925
Source: JFWP
Date Processed by STIC: 12/20/2005

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number:

10/559,925

CRF Edit Date:

12/21/2005

Edited by:

DA

— Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

— Corrected the SEQ ID NO. Sequence numbers edited were:

— Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

~~Deleted: _____ invalid beginning/end-of-file text ; _____ page numbers~~

— Inserted mandatory headings/numeric identifiers, specifically:

— Moved responses to same line as heading/numeric identifier, specifically:

— Other:



IFWO

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/559,925

DATE: 12/21/2005
TIME: 10:43:53

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\12202005\J559925.raw

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3 <110> APPLICANT: Yacoby-Zeevi, Oron
4     Peretz, Tuvia
5     Miron, Daphna
6     Shlomi, Yinon
7     Pecker, Iris
8     Ayal-Hershkovitz, Maty
9     Feinstein, Elena
10    Van Gelder, Joel M.
11    Vlodavsky, Israel
12    Friedmann, Yael
14 <120> TITLE OF INVENTION: HEPARANASE ACTIVITY NEUTRALIZING ANTI- HEPARANASE MONOCLONAL
15 ANTIBODY AND OTHER ANTI-HEPARANASE ANTIBODIES
17 <130> FILE REFERENCE: 30337
C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/559,925
C--> 19 <141> CURRENT FILING DATE: 2005-12-08
19 <160> NUMBER OF SEQ ID NOS: 11
21 <170> SOFTWARE: PatentIn version 3.2
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 386
25 <212> TYPE: PRT
26 <213> ORGANISM: Homo sapiens
29 <220> FEATURE:
30 <221> NAME/KEY: misc_feature
31 <223> OTHER INFORMATION: 45 kDa subunit of mature processed heparanase dimer
33 <400> SEQUENCE: 1
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39 Tyr Thr Phe Ala Asn Cys Ser Gly Leu Asp Leu Ile Phe Gly Leu Asn
40          20          25           30
43 Ala Leu Leu Arg Thr Ala Asp Leu Gln Trp Asn Ser Ser Asn Ala Gln
44          35          40           45
47 Leu Leu Leu Asp Tyr Cys Ser Ser Lys Gly Tyr Asn Ile Ser Trp Glu
48          50          55           60
51 Leu Gly Asn Glu Pro Asn Ser Phe Leu Lys Lys Ala Asp Ile Phe Ile
52 65          70          75           80
55 Asn Gly Ser Gln Leu Gly Glu Asp Phe Ile Gln Leu His Lys Leu Leu
56          85          90           95
59 Arg Lys Ser Thr Phe Lys Asn Ala Lys Leu Tyr Gly Pro Asp Val Gly
60          100         105          110
63 Gln Pro Arg Arg Lys Thr Ala Lys Met Leu Lys Ser Phe Leu Lys Ala
64          115         120          125
67 Gly Gly Glu Val Ile Asp Ser Val Thr Trp His His Tyr Tyr Leu Asn
68          130         135          140

```

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71 Gly Arg Thr Ala Thr Arg Glu Asp Phe Leu Asn Pro Asp Val Leu Asp
72 145 150 155 160
75 Ile Phe Ile Ser Ser Val Gln Lys Val Phe Gln Val Val Glu Ser Thr
76 165 170 175
79 Arg Pro Gly Lys Lys Val Trp Leu Gly Glu Thr Ser Ser Ala Tyr Gly
80 180 185 190
83 Gly Gly Ala Pro Leu Leu Ser Asp Thr Phe Ala Ala Gly Phe Met Trp
84 195 200 205
87 Leu Asp Lys Leu Gly Leu Ser Ala Arg Met Gly Ile Glu Val Val Met
88 210 215 220
91 Arg Gln Val Phe Phe Gly Ala Gly Asn Tyr His Leu Val Asp Glu Asn
92 225 230 235 240
95 Phe Asp Pro Leu Pro Asp Tyr Trp Leu Ser Leu Leu Phe Lys Lys Leu
96 245 250 255
99 Val Gly Thr Lys Val Leu Met Ala Ser Val Gln Gly Ser Lys Arg Arg
100 260 265 270
103 Lys Leu Arg Val Tyr Leu His Cys Thr Asn Thr Asp Asn Pro Arg Tyr
104 275 280 285
107 Lys Glu Gly Asp Leu Thr Leu Tyr Ala Ile Asn Leu His Asn Val Thr
108 290 295 300
111 Lys Tyr Leu Arg Leu Pro Tyr Pro Phe Ser Asn Lys Gln Val Asp Lys
112 305 310 315 320
115 Tyr Leu Leu Arg Pro Leu Gly Pro His Gly Leu Leu Ser Lys Ser Val
116 325 330 335
119 Gln Leu Asn Gly Leu Thr Leu Lys Met Val Asp Asp Gln Thr Leu Pro
120 340 345 350
123 Pro Leu Met Glu Lys Pro Leu Arg Pro Gly Ser Ser Leu Gly Leu Pro
124 355 360 365
127 Ala Phe Ser Tyr Ser Phe Phe Val Ile Arg Asn Ala Lys Val Ala Ala
128 370 375 380
131 Cys Ile
132 385
135 <210> SEQ ID NO: 2
136 <211> LENGTH: 535
137 <212> TYPE: PRT
138 <213> ORGANISM: Mus musculus
140 <400> SEQUENCE: 2
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146 Ala Gln Gly Ala Pro Ala Gly Thr Ala Pro Thr Asp Asp Val Val Asp
147 20 25 30
150 Leu Glu Phe Tyr Thr Lys Arg Pro Leu Arg Ser Val Ser Pro Ser Phe
151 35 40 45
154 Leu Ser Ile Thr Ile Asp Ala Ser Leu Ala Thr Asp Pro Arg Phe Leu
155 50 55 60
158 Thr Phe Leu Gly Ser Pro Arg Leu Arg Ala Leu Ala Arg Gly Leu Ser
159 65 70 75 80
162 Pro Ala Tyr Leu Arg Phe Gly Gly Thr Lys Thr Asp Phe Leu Ile Phe
163 85 90 95

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Input Set : A:\pto.da.txt
Output Set: N:\CRF4\12202005\J559925.raw

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166 Asp Pro Asp Lys Glu Pro Thr Ser Glu Glu Arg Ser Tyr Trp Lys Ser
167      100          105          110
170 Gln Val Asn His Asp Ile Cys Arg Ser Glu Pro Val Ser Ala Ala Val
171      115          120          125
174 Leu Arg Lys Leu Gln Val Glu Trp Pro Phe Gln Glu Leu Leu Leu Leu
175      130          135          140
178 Arg Glu Gln Tyr Gln Lys Glu Phe Lys Asn Ser Thr Tyr Ser Arg Ser
179 145          150          155          160
182 Ser Val Asp Met Leu Tyr Ser Phe Ala Lys Cys Ser Gly Leu Asp Leu
183          165          170          175
186 Ile Phe Gly Leu Asn Ala Leu Leu Arg Thr Pro Asp Leu Arg Trp Asn
187          180          185          190
190 Ser Ser Asn Ala Gln Leu Leu Leu Asp Tyr Cys Ser Ser Lys Gly Tyr
191          195          200          205
194 Asn Ile Ser Trp Glu Leu Gly Asn Glu Pro Asn Ser Phe Trp Lys Lys
195 210          215          220
198 Ala His Ile Leu Ile Asp Gly Leu Gln Leu Gly Glu Asp Phe Val Glu
199 225          230          235          240
202 Leu His Lys Leu Leu Gln Arg Ser Ala Phe Gln Asn Ala Lys Leu Tyr
203          245          250          255
206 Gly Pro Asp Ile Gly Gln Pro Arg Gly Lys Thr Val Lys Leu Leu Arg
207 260          265          270
210 Ser Phe Leu Lys Ala Gly Gly Glu Val Ile Asp Ser Leu Thr Trp His
211          275          280          285
214 His Tyr Tyr Leu Asn Gly Arg Ile Ala Thr Lys Glu Asp Phe Leu Ser
215 290          295          300
218 Ser Asp Ala Leu Asp Thr Phe Ile Leu Ser Val Gln Lys Ile Leu Lys
219 305          310          315          320
222 Val Thr Lys Glu Ile Thr Pro Gly Lys Lys Val Trp Leu Gly Glu Thr
223          325          330          335
226 Ser Ser Ala Tyr Gly Gly Ala Pro Leu Leu Ser Asn Thr Phe Ala
227          340          345          350
230 Ala Gly Phe Met Trp Leu Asp Lys Leu Gly Leu Ser Ala Gln Met Gly
231          355          360          365
234 Ile Glu Val Val Met Arg Gln Val Phe Phe Gly Ala Gly Asn Tyr His
235 370          375          380
238 Leu Val Asp Glu Asn Phe Glu Pro Leu Pro Asp Tyr Trp Leu Ser Leu
239 385          390          395          400
242 Leu Phe Lys Lys Leu Val Gly Pro Arg Val Leu Leu Ser Arg Val Lys
243          405          410          415
246 Gly Pro Asp Arg Ser Lys Leu Arg Val Tyr Leu His Cys Thr Asn Val
247          420          425          430
250 Tyr His Pro Arg Tyr Gln Glu Gly Asp Leu Thr Leu Tyr Val Leu Asn
251          435          440          445
254 Leu His Asn Val Thr Lys His Leu Lys Val Pro Pro Pro Leu Phe Arg
255          450          455          460
258 Lys Pro Val Asp Thr Tyr Leu Leu Lys Pro Ser Gly Pro Asp Gly Leu
259 465          470          475          480
262 Leu Ser Lys Ser Val Gln Leu Asn Gly Gln Ile Leu Lys Met Val Asp

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Input Set : A:\pto.da.txt
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263	485	490	495
266	Glu Gln Thr Leu Pro Ala Leu Thr Glu Lys Pro Leu Pro Ala Gly Ser		
267	500	505	510
270	Ala Leu Ser Leu Pro Ala Phe Ser Tyr Gly Phe Phe Val Ile Arg Asn		
271	515	520	525
274	Ala Lys Ile Ala Ala Cys Ile		
275	530	535	
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279	<211> LENGTH: 536		
280	<212> TYPE: PRT		
281	<213> ORGANISM: Rattus norvegicus		
283	<400> SEQUENCE: 3		
285	Met Leu Arg Pro Leu Leu Leu Trp Leu Trp Gly Arg Leu Arg Ala		
286	1 5	10	15
289	Leu Thr Gln Gly Thr Pro Ala Gly Thr Ala Pro Thr Lys Asp Val Val		
290	20 25	30	
293	Asp Leu Glu Phe Tyr Thr Lys Arg Leu Phe Gln Ser Val Ser Pro Ser		
294	35 40	45	
297	Phe Leu Ser Ile Thr Ile Asp Ala Ser Leu Ala Thr Asp Pro Arg Phe		
298	50 55	60	
301	Leu Thr Phe Leu Gly Ser Pro Arg Leu Arg Ala Leu Ala Arg Gly Leu		
302	65 70	75	80
305	Ser Pro Ala Tyr Leu Arg Phe Gly Gly Thr Lys Thr Asp Phe Leu Ile		
306	85 90	95	
309	Phe Asp Pro Asn Lys Glu Pro Thr Ser Glu Glu Arg Ser Tyr Trp Gln		
310	100 105	110	
313	Ser Gln Asp Asn Asn Asp Ile Cys Gly Ser Glu Arg Val Ser Ala Asp		
314	115 120	125	
317	Val Leu Arg Lys Leu Gln Met Glu Trp Pro Phe Gln Glu Leu Leu Leu		
318	130 135	140	
321	Leu Arg Glu Gln Tyr Gln Arg Glu Phe Lys Asn Ser Thr Tyr Ser Arg		
322	145 150	155	160
325	Ser Ser Val Asp Met Leu Tyr Ser Phe Ala Lys Cys Ser Arg Leu Asp		
326	165 170	175	
329	Leu Ile Phe Gly Leu Asn Ala Leu Leu Arg Thr Pro Asp Leu Arg Trp		
330	180 185	190	
333	Asn Ser Ser Asn Ala Gln Leu Leu Asn Tyr Cys Ser Ser Lys Gly		
334	195 200	205	
337	Tyr Asn Ile Ser Trp Glu Leu Gly Asn Glu Pro Asn Ser Phe Trp Lys		
338	210 215	220	
341	Lys Ala Gln Ile Ser Ile Asp Gly Leu Gln Leu Gly Glu Asp Phe Val		
342	225 230	235	240
345	Glu Leu His Lys Leu Leu Gln Lys Ser Ala Phe Gln Asn Ala Lys Leu		
346	245 250	255	
349	Tyr Gly Pro Asp Ile Gly Gln Pro Arg Gly Lys Thr Val Lys Leu Leu		
350	260 265	270	
353	Arg Ser Phe Leu Lys Ala Gly Gly Glu Val Ile Asp Ser Leu Thr Trp		
354	275 280	285	
357	His His Tyr Tyr Leu Asn Gly Arg Val Ala Thr Lys Glu Asp Phe Leu		

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Input Set : A:\pto.da.txt
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358	290	295	300
361	Ser Ser Asp Val Leu Asp Thr Phe Ile Leu Ser Val Gln Lys Ile Leu		
362	305	310	315
365	Lys Val Thr Lys Glu Met Thr Pro Gly Lys Lys Val Trp Leu Gly Glu		320
366	325	330	335
369	Thr Ser Ser Ala Tyr Gly Gly Ala Pro Leu Leu Ser Asn Thr Phe		
370	340	345	350
373	Ala Ala Gly Phe Met Trp Leu Asp Lys Leu Gly Leu Ser Ala Gln Leu		
374	355	360	365
377	Gly Ile Glu Val Val Met Arg Gln Val Phe Phe Gly Ala Gly Asn Tyr		
378	370	375	380
381	His Leu Val Asp Glu Asn Phe Glu Pro Leu Pro Asp Tyr Trp Leu Ser		
382	385	390	395
385	Leu Leu Phe Lys Lys Leu Val Gly Pro Lys Val Leu Met Ser Arg Val		400
386	405	410	415
389	Lys Gly Pro Asp Arg Ser Lys Leu Arg Val Tyr Leu His Cys Thr Asn		
390	420	425	430
393	Val Tyr His Pro Arg Tyr Arg Glu Gly Asp Leu Thr Leu Tyr Val Leu		
394	435	440	445
397	Asn Leu His Asn Val Thr Lys His Leu Lys Leu Pro Pro Pro Met Phe		
398	450	455	460
401	Ser Arg Pro Val Asp Lys Tyr Leu Leu Lys Pro Phe Gly Ser Asp Gly		
402	465	470	475
405	Leu Leu Ser Lys Ser Val Gln Leu Asn Gly Gln Thr Leu Lys Met Val		480
406	485	490	495
409	Asp Glu Gln Thr Leu Pro Ala Leu Thr Glu Lys Pro Leu Pro Ala Gly		
410	500	505	510
413	Ser Ser Leu Ser Val Pro Ala Phe Ser Tyr Gly Phe Phe Val Ile Arg		
414	515	520	525
417	Asn Ala Lys Ile Ala Ala Cys Ile		
418	530	535	
421	<210> SEQ ID NO: 4		
422	<211> LENGTH: 543		
423	<212> TYPE: PRT		
424	<213> ORGANISM: Homo sapiens		
426	<400> SEQUENCE: 4		
428	Met Leu Leu Arg Ser Lys Pro Ala Leu Pro Pro Pro Leu Met Leu Leu		
429	1	5	10
432	15		
433	Leu Leu Gly Pro Leu Gly Pro Leu Ser Pro Gly Ala Leu Pro Arg Pro		
436	20	25	30
437	Ala Gln Ala Gln Asp Val Val Asp Leu Asp Phe Phe Thr Gln Glu Pro		
440	35	40	45
441	Leu His Leu Val Ser Pro Ser Phe Leu Ser Val Thr Ile Asp Ala Asn		
444	50	55	60
445	Leu Ala Thr Asp Pro Arg Phe Leu Ile Leu Leu Gly Ser Pro Lys Leu		
448	65	70	75
449	80		
452	Arg Thr Leu Ala Arg Gly Leu Ser Pro Ala Tyr Leu Arg Phe Gly Gly		
	85	90	95
	Thr Lys Thr Asp Phe Leu Ile Phe Asp Pro Lys Lys Glu Ser Thr Phe		

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/559,925

DATE: 12/21/2005

TIME: 10:43:54

Input Set : A:\pto.da.txt

Output Set: N:\CRF4\12202005\J559925.raw

L:19 M:270 C: Current Application Number differs, Replaced Current Application No

L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date

**Raw Sequence Listing before editing,
for reference only**



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/559,925

DATE: 12/20/2005
TIME: 10:26:33

Input Set : A:\30337 Sequence Listing.txt
Output Set: N:\CRF4\12202005\J559925.raw

3 <110> APPLICANT: Yacoby-Zeevi, Oron
 4 Peretz, Tuvia
 5 Miron, Daphna
 6 Shlomi, Yinon
 7 Pecker, Iris
 8 Ayal-Hershkovitz, Maty
 9 Feinstein, Elena
 10 Van Gelder, Joel M.
 11 Vlodavsky, Israel
 12 Friedmann, Yael
 14 <120> TITLE OF INVENTION: HEPARANASE ACTIVITY NEUTRALIZING ANTI- HEPARANASE MONOCLONAL
 15 ANTIBODY AND OTHER ANTI-HEPARANASE ANTIBODIES
 17 <130> FILE REFERENCE: 30337
 C--> 19 <140> CURRENT APPLICATION NUMBER: US/10/559,925
 C--> 19 <141> CURRENT FILING DATE: 2005-12-08
 19 <160> NUMBER OF SEQ ID NOS: 11
 21 <170> SOFTWARE: PatentIn version 3.2

Does Not Comply
 Corrected Diskette Needed
 (CPR)

ERRORED SEQUENCES

777 <210> SEQ ID NO: 11
 778 <211> LENGTH: 74
 779 <212> TYPE: PRT
 780 <213> ORGANISM: Homo sapiens
 783 <220> FEATURE:
 784 <221> NAME/KEY: misc_feature
 785 <223> OTHER INFORMATION: 8 kDa subunit of mature processed heparanase dimer
 787 <400> SEQUENCE: 11
 789 Gln Asp Val Val Asp Leu Asp Phe Phe Thr Gln Glu Pro Leu His Leu
 790 1 5 10 15
 793 Val Ser Pro Ser Phe Leu Ser Val Thr Ile Asp Ala Asn Leu Ala Thr
 794 20 25 30
 797 Asp Pro Arg Phe Leu Ile Leu Gly Ser Pro Lys Leu Arg Thr Leu
 798 35 40 45
 801 Ala Arg Gly Leu Ser Pro Ala Tyr Leu Arg Phe Gly Gly Thr Lys Thr
 802 50 55 60
 805 Asp Phe Leu Ile Phe Asp Pro Lys Lys Glu
 806 65 70
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/559,925

DATE: 12/20/2005

TIME: 10:26:35

Input Set : A:\30337 Sequence Listing.txt

Output Set: N:\CRF4\12202005\J559925.raw

L:19 M:270 C: Current Application Number differs, Replaced Current Application No

L:19 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:811 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11